

## SEQUENCE LISTING JC10 Recd PCT/EP 2.0 MAR 2002

<110> Fandke, Markus  
Gasch, Alexander  
Berghof, Kornelia

<120> Method and nucleic acids for the detection of microorganisms  
relevant to brewing

<130> 216087

<140>

<141> 2002-03-20

<150> PCT/EP00/08808

<151> 2000-09-08

<150> DE 199 45 964.9

<151> 1999-09-24

<160> 107

<170> PatentIn Ver. 2.1

<210> 1

<211> 267

<212> DNA

<213> Lactobacillus brevis

<400> 1

|   |     |
|---|-----|
| tatatggaag taagaccct gagagatgat caggtagata ggctggaagt agcagcgccg  | 60  |
| tgaggcgtgg agcggaccag tactaatcgg tcgaggactt aaccaagtca acaacgtagt | 120 |
| tgtttcgaga ataattgaat aatatctagt tttagaggaa gaagttctct tatagtgtgg | 180 |
| tggcgatagc ctgaaggata cacctgttcc catgccgaac acagaagtta agcttcagca | 240 |
| cgccgatagt agttggggga tcgcccc                                     | 267 |

<210> 2

<211> 326

<212> DNA

<213> Lactobacillus lindneri

<400> 2

|   |     |
|---|-----|
| ccattcctat atggaagtaa gactcctgaa agatgatcag gtcgataggt tagaagtgga | 60  |
| agcatagtga tatgtgaagc ggactaatac taatcagtcg aggacttaac caaggaagac | 120 |
| acagggttaa atcaaagttg aacagagaag atattatcta gttttgagag aacgaagttc | 180 |
| gctcaggctt atgaaaaata agcatagtgt ggtggcgata gcctgaagga tacacctgtt | 240 |
| cccatgccga acacagaagt taagcttcag caccgcaaaa gtagttgggg gatcgcccc  | 300 |
| tgcgaggata ggacgatggt catagc                                      | 326 |

<210> 3

<211> 351

<212> DNA

<213> Lactobacillus casei

<400> 3

|   |     |
|---|-----|
| ccattcctat atggaagtaa gaccctgag agatgatcag gtagataggc tggaagtgga  | 60  |
| agtgcagcga tgcattgagc ggaccagtac taatcggtcg aggacttaac caagtagagc | 120 |
| gtgagcagga gcgcttagaa accggagcat aagcgggcct gagttcggtg gccgggtttt | 180 |
| ggccaatgga ttcagggttc ttatgtggag gtttctgcga ctgcgaacgc gtttcgatga | 240 |

aatacactgg ttcccgacaa cacaaaaaca acaatgatag ccagttttga gagcgcaaag 300  
 ttctcataag tgtggtggcg atagcaagaa ggatacacct gttcccatgc c 351

<210> 4

<211> 414

<212> DNA

<213> *Lactobacillus paracasei*

<400> 4

ccattcctat atggaagtaa gacccttgag agatgatcag gtagataggc tggaagtgga 60  
 agtgcagcga tgcattggagc ggaccagtac taatcggtcg aggacttaac caagtaagag 120  
 tgtgagcagg agcgggttaga aaccggagca taagcgggcc tgagcgtgat ggccgggctt 180  
 tggccattgc ggtcagggtc cttatgtgca ggtttctgcg actgcgaaca cgtttcgatg 240  
 acaagtacgt taagttcaag gcagcaatta aacaatgata gctagttttg agagcgcaaa 300  
 gttctcataa gtgtggtggc gatagcaaga aggatacacc tgttcccatg ccgaacacag 360  
 aagttaagct tcttcacgcc gagagtagtt ggtgggaaac tgctgcgag gata 414

<210> 5

<211> 338

<212> DNA

<213> *Lactobacillus paracasei*

<400> 5

ccattcctat atggaagtaa gacccttgag agatgatcag gtagataggc tggaagtgga 60  
 agtgcagcga tgcattggagc ggaccagtac taatcggtcg aggacttaac caagtaagcg 120  
 tgcaagcagg agcaggtttc tgcgactgcg aacacatttc gatgacaagt acgttaagtt 180  
 caaggcagca attaaacgat gatagccagt tttgagagcg caaagttctc ataagtgtgg 240  
 tggcgatagc aagaaggata cacctgttcc catgccgaac acagaagtta agcttcttca 300  
 cgccgagagt agttggtggg aaactgcctg cgaggata 338

<210> 6

<211> 317

<212> DNA

<213> *Lactobacillus coryniformis* ssp. *coryniformis*

<400> 6

ctcgagttga gatttcccat tcctttatgg aagtaagacc cctgagagat gatcaggtag 60  
 ataggttgga agtggacgtg ccgtgaggca tggagcggac caataactaat cggtcgagga 120  
 cttaaccaag tagcatgtac gtagtgtagg ttttaaggga aagaaatgaa tatccagttt 180  
 tgagagcgca acgttctcag aaagtgggtg ggtggcgata gcaagaagga tacacctgtt 240  
 cccatgtcga acacagaagt taagcttctt agcgccgaga gtagttgggg gagcaccccc 300  
 tgcgaggata ggacgat 317

<210> 7

<211> 317

<212> DNA

<213> *Lactobacillus coryniformis* ssp. *torquens*

<400> 7

ctcgagatga gatttcccat tcctttatgg aagtaagacc cctgagagat gatcaggtag 60  
 ataggttgga agtggacgtg ccgtgaggca tggagcggac caataactaat cggtcgagga 120  
 cttaaccaag tagcatgtac gtggtgtagg ttttaaggga aagaaatgaa tatccagttt 180  
 tgagagcgca acgttctcag aaagtgggtg ggtggcgata gcaagaagga tacacctgtt 240  
 cccatgtcga acacagaagt taagcttctt agcgccgaga gtagttgggg gagcaccccc 300  
 tgcgaggata ggacgat 317

<210> 8  
 <211> 336  
 <212> DNA  
 <213> *Lactobacillus curvatus*

<400> 8  
 acgcctcgag atgagatttc ccattccttt atggaagtaa gacccctgaa agatgatcag 60  
 gtagataggc taggagtggg agtacagcga tgtatggagc ggactagtag taatcggtag 120  
 aggacttaac caaagggtgca atgttaggct tttgaaatga aatattactt attatgcagt 180  
 tttgagagaa cgaagttctt ctgagtgcgc aagcacaaaa tagtgtgggtg gcgatagcaa 240  
 gaaggataca cctgttccca tgtcgaacac agaagttaag cttcttagcg ccgatagtag 300  
 ttggtgggaa actacctgcg aggataggac gatggg 336

<210> 9  
 <211> 335  
 <212> DNA  
 <213> *Pediococcus damnosus*

<400> 9  
 gatgagattt cccattccat ttatggaagt aagaccctg agagatgatc aggtagatag 60  
 gttgggagtg gaagtgtagt gatacatgga gcggaccaat actaatcggg cgaggactta 120  
 accacaaagt ggtgttctca agagaaggat tcgatattat ttagttttga gagaataaat 180  
 ttctttcaca cgagccgcgt aagtggatcg gagaagtgtg gtgacgatag tgagaaggat 240  
 acacctgttc ccatgtcgaa cacagaagtt aagcttctta acgccgagag tagttggggg 300  
 atcgctccct gcgaggatag gacgatgggc aatag 335

<210> 10  
 <211> 326  
 <212> DNA  
 <213> *Pediococcus inopinatus*

<400> 10  
 agatgagatt tcccattcca tttatggaag taagaccctt gagagatgat caggtagata 60  
 ggttgggagt ggaagtgtag tgatacatgg agcggaccaa tactaatcgg tcgaggactt 120  
 aaccacaaag tgggtgttctc aaagagaaga tttcgatatt atttagtttt gagagaataa 180  
 atttctttca cagcagccgc ggaagtggat cggagaagtg tgggtgacgat agtgagaagg 240  
 atacacctgt tcccatgtcg aacacagaag ttaagcttct taacgccgag agtagttggg 300  
 ggatcgctcc ctgcgaggat aggacg 326

<210> 11  
 <211> 403  
 <212> DNA  
 <213> *Pectinatus cerevisiiphilus*

<400> 11  
 aagtgcctgaa agcatctaag cgtgaaacct gccttaagat gaggtttccc agagccgtaa 60  
 ggcttggaag gcaccttgaa taagacgagg tagataggcc gggagttagaa gtacagtaat 120  
 gtacgaagcg gactggtact aataagccga gagcttaact taaaatcatc gaaaaaatg 180  
 tttggtctga gatttcttct gtgaagtttt gagtgtgcaa gacactctgg ttgaagggca 240  
 gggaacgtga gagcgtaaaa ctgcggactt tggctcaaag agttaagca tctggtgacg 300  
 atacctggat ggatccacct gttcccatto cgaacacagt agttaagcat ccacaggctg 360  
 aaggtacttg gggggcgacc ccctgggaaa ataggacact gcc 403

<210> 12  
 <211> 434  
 <212> DNA  
 <213> *Pectinatus frisingensis*

<400> 12  
aagtgcgtgaa agcatcctaag cgtgaaacca gctttaagat gaggtttccc agaacgcaag 60  
tttgggaaggc accttgaaga agacgaggta gataggccgg gagtggaagt atggtgacat 120  
atgaagcggga ctggtactaa taagccgaga gcttaacttg atttcatcaa aaaagagaaa 180  
tgtttggtca gagattttct tctgtgaagt tttgagtgtg caagaacact cgagagtata 240  
taggtaaagg aaaagcagca gataagtttc ctggttactg tatataccgg ctgaggtgct 300  
gaggcactga aggccagaac atctggtggc gatacctgga tggatccacc tgttcccatt 360  
ccgaacacag tagttaagca tccacaggcc gaaggtactt ggggggcagc cccctgcgaa 420  
aataggacac cgcc 434

<210> 13

<211> 641

<212> DNA

<213> *Pectinatus spec.* DSM20764

<400> 13  
aagtgcgtgaa agcatcctaag cgtgaaacct gccttaagat gaggtttccc agagccgtaa 60  
ggcttggaag gcaccttgaa gatgacgagg tagataggcc gggagtagaa gtatggtgac 120  
atacgaagcg gactggtact aataagccga gagcttaact taatttcac tataaatgtt 180  
tggctcctgat ttcttctgtg aagttttgag tgtgcaagat cactcatgaa agtatatagg 240  
taaaggggaaa gcagcagatt agttcctggg ttactttata tatgagcact aaggtgcaga 300  
aaagaacggt tgaggaaacg cggcggttct aaactcact tgcgtgctga ttatctcaat 360  
gctaaagcat taagataatt ttagaggaaa cgcgcgttca ctacggttca ctctgcgtac 420  
tttatttcta agtgctgaag cactaagaag ggcaaggaaa cgcgtcgttc gcgatgctca 480  
ctttgcgtac ttcatctcta gactgctaaa gcagtaagat ctgaagcatc tgggtggcgat 540  
acctggatgg atccacctgt tcccattccg aacacagtag ttaagcatcc acaggccgaa 600  
ggtacttggg gggcagcccc ctgcgagagt aggacatcgc c 641

<210> 14

<211> 495

<212> DNA

<213> *Pectinatus spec.* DSM20764

<400> 14  
aagtgcgtgaa agcatcctaag cgtgaaacct gccttaagat gaggtttccc agagccgtaa 60  
ggcttggaag gcaccttgaa gatgacgagg tagataggcc gggagtagaa gtatggtgac 120  
atacgaagcg gactggtact aataagccga gagcttaact taatttcac tataaatgtt 180  
tggctcctgat ttcttctgtg aagttttgag tgtgcaagat cactcatgaa agtatatagg 240  
taaaggggaaa gcagattagt tcctggttta ctttatatat gagcactaag gtgcagaaaa 300  
gaacgtctaa ggaaacgcgg cgttcgtagg ctactctgc gtacttcac tctagactgc 360  
taaagcagta agatctgaag catctggtgg cgatacctgg atggatccac ctgttcccat 420  
tccgaacaca gtagttaagc atccacaggc cgaaggtact tggggggcag cccctgcgaa 480  
aagtaggaca ccgcc 495

<210> 15

<211> 546

<212> DNA

<213> *Megasphaera cerevisiae*

<400> 15  
gcatcctaagc gtgaaaccag cctagagatg aggtttctca ttacgaaagt aagtaaggtc 60  
ccatgaagac gacatggtag ataggccggg agtggacgta cagtaatgta tggagcggac 120  
cggactaat agaccgagga cttgacttaa gcagggaacc cattttaaag aagcgaagcg 180  
acgcataaaa tggagtgaat cgcttatacc gaatcgaga ttcggtaaaag cagcggagaa 240  
taccaatgca gcggcaacac cagttagcat aaactaagcg gattcggagt ggggtgaggga 300  
gtttcgtagc agcgtaggct aacccaacca ccgctttcga agaaggcgaa tggtttgaaa 360  
aagagtacat gcgaagaaac gacgaaagac tcacaaccaa aacatacaaa ctaagtagat 420

gacattagag tcacaccgat tgttaagatc cgaaatactt ttcgatgtag ttgtcaggat 480  
 acgaatcctg aaacgaattc agtgggtgatg gctgcaggga tccacctgtt cccataccga 540  
 acacag 546

<210> 16  
 <211> 306  
 <212> DNA  
 <213> *Megasphaera cerevisiae*

<400> 16  
 gcatctaacc gtgaaaccag cctagagatg aggtttctca ttacgaaagt aagtaaggtc 60  
 ccatgaagac gacatggtag ataggccggg agtggacgta cagtaaatgta tggagcggac 120  
 cggtaactaat agaccgagga cttgacttaa gcaaagaagc aatagaaaga accatgtaga 180  
 tgggtgtaaga gttagacggg tagttaaggc ccgaaatact tttcgatgta gttgtcagga 240  
 tacgaatcct gaaacgaatt cagtgggtgat ggctgcaggg accacctgtt cccataccga 300  
 acacag 306

<210> 17  
 <211> 449  
 <212> DNA  
 <213> *Selenomonas lacticifex*

<400> 17  
 aagtgtctgaa agcatctagg cgtgaagcct gtcccgagat gaagtatctc atggagtaat 60  
 ccagtaagat tccttgaaga agacaaggta gatagggttg gagtgtaagc atcgtaaggc 120  
 gttcagcggg ccaataactaa taaatcgagg gcttaacttt acagacctgt ccaagaagcg 180  
 aagcggattg ggtaacaggc cgtatgcgaa aacatcccaa gaatcgagtc cgaagggcga 240  
 agatgattgg cagatgttga ccgctaataa tctagaatgt ttcgatacaa tttttcttct 300  
 gtatagtttt gagtggacat cgttcattca ataatatcca gtgacgatag ctgagtggtg 360  
 ccacctgttc ccataccgaa cacagtagtt aagcactcat acgccgaaag tacttgtctg 420  
 gaaacgggct gcgagaatag gacgtcgcc 449

<210> 18  
 <211> 343  
 <212> DNA  
 <213> *Selenomonas lacticifex*

<400> 18  
 aagtgtctgaa agcatctaag cgtgaagcct gtcccgagat gaagtatctc atggagtaat 60  
 ccagtaagat tccttgaaga agacaaggta gatagggttg gagtgtaagc atcgtaaggc 120  
 gttcagcggg ccaataactaa taaatcgagg gcttatctta ataacttaga atgtttcgat 180  
 acaatttttc ttctgtatag ttttgagtgg acatgggttca ttcaataata tccagtgcag 240  
 atagctgagt ggtaccacct gttcccatac cgaacacagt agttaagcac tcatacgccg 300  
 aaagtacttg tctggaaacg ggctgcgaaa ataggacgcc gcc 343

<210> 19  
 <211> 395  
 <212> DNA  
 <213> *Zymophilus raffinivorans*

<400> 19  
 aagtgtctgaa agcatctaag cgtgaaacca gccttaagat gaggtttctc acagagcaat 60  
 ctggtaagac cccttgaaga agacaaggta gatagggtcg gagtggaagc gcagtaaatg 120  
 gtgcagcggg ccgatactaa taggtcgagg gcttgactta aagccagaac gaaaactaaa 180  
 atgcgaacat ttctttcttc tgtatagttt tgagagaaca aactcttaag atggagtagt 240  
 ctgaggcgaa agcggaaaggc agcgatatct aaaaaaagaa tatctggtag tgatagccaa 300  
 gtggaccac ctgttcccat accgaacaca gtagttaagc acttgaacgt cgaaagtact 360

tgggtggaaa cgccctgcga aaataggaca ccgcc

395

<210> 20  
 <211> 395  
 <212> DNA  
 <213> *Zymophilus paucivorans*

<400> 20  
 aagtgtctgaa agcatctaa cgtgaaacca gccttaagat gaggtttctc acagagcaat 60  
 ctggtgaagac cccttgaaga agacaaggta gataggtcgg gagtgggaagc gcagtaatgt 120  
 gtgtagcggg ccgatactaa taggtcgagg gcttgactta aagccagaac gaattctaaa 180  
 atgcgaacat ttctttcttc tgtatagttt tgagagaaca gactcttaag atgagcagtc 240  
 tgaggcgaaa gctaaaggca gcgatattcta aaaaaaagaa tatctggtag tgatagccaa 300  
 gtggaccac ctgttcccat accgaacaca gtagttaagc acttgaacgt cgaaagtact 360  
 tgggtggaaa cgccctggga aaataggaca ccgcc 395

<210> 21  
 <211> 21  
 <212> DNA  
 <213> Artificial sequence

<220>  
 <223> Description of the artificial sequence: specific  
 sequence for *Lactobacillus brevis*

<400> 21  
 ccaagtcaac aacgtagttg t 21

<210> 22  
 <211> 23  
 <212> DNA  
 <213> Artificial sequence

<220>  
 <223> Description of the artificial sequence: specific  
 sequence for *Lactobacillus lindneri*

<400> 22  
 gacacagggt taaatcaaag ttg 23

<210> 23  
 <211> 20  
 <212> DNA  
 <213> Artificial sequence

<220>  
 <223> Description of the artificial sequence: specific  
 sequence for *Lactobacillus casei* and *Lactobacillus*  
*paracasei*

<400> 23  
 aggtttctgc gactgcgaac 20

<210> 24  
 <211> 25  
 <212> DNA

<213> Artificial sequence

<220>

<223> Description of the artificial sequence: specific  
sequence for *Lactobacillus coryniformis*

<400> 24

atgtacgtag tgtagttta agggc

25

<210> 25

<211> 20

<212> DNA

<213> Artificial sequence

<220>

<223> Description of the artificial sequence: specific  
sequence for *Lactobacillus curvatus*

<400> 25

cttctcagtg cgcaagcaca

20

<210> 26

<211> 22

<212> DNA

<213> Artificial sequence

<220>

<223> Description of the artificial sequence: specific  
sequence for *Pediococcus damnosus*

<400> 26

gtgttctcaa gagaaggatt cg

22

<210> 27

<211> 27

<212> DNA

<213> Artificial sequence

<220>

<223> Description of the artificial sequence: specific  
sequence for *Pediococcus inopinatus*

<400> 27

gttctcaaag agaagatttc gatatta

27

<210> 28

<211> 23

<212> DNA

<213> Artificial sequence

<220>

<223> Description of the artificial sequence: specific  
sequence for *Pectinatus cerevisiophilus*

<400> 28

tgagagcgta aaactgcgga ctt

23

<210> 29  
 <211> 22  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: specific  
 sequence for *Pectinatus frisingensis*

<400> 29  
 cagataagtt tcctggttac tg 22

<210> 30  
 <211> 23  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: specific  
 sequence for *Pectinatus spec.* DSM 20764

<400> 30  
 cactaaggtg cagaaaagaa cgt 23

<210> 31  
 <211> 26  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: specific  
 sequence for *Megasphaera cerevisiae*

<400> 31  
 cttttcgatg tagttgtcag gatacg 26

<210> 32  
 <211> 25  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: specific  
 sequence for *Selenomonas lacticifex*

<400> 32  
 gttcattcaa taatatccag tgacg 25

<210> 33  
 <211> 23  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: specific  
 sequence for *Zymophilus raffinovorans*



<400> 33  
aactcttaag atggagyagt ctg 23

<210> 34  
<211> 22  
<212> DNA  
<213> Artifical sequence

<220>  
<223> Description of the artifical sequence: specific  
sequence for Zymophilus paucivorans

<400> 34  
actcttaaga tgagcagtct ga 22

<210> 35  
<211> 21  
<212> DNA  
<213> Artifical sequence

<220>  
<223> Description of the artifical sequence: specific  
sequence for the genus Pediococcus

<400> 35  
agtstagtga tacatggagc g 21

<210> 36  
<211> 22  
<212> DNA  
<213> Artifical sequence

<220>  
<223> Description of the artifical sequence: specific  
sequence for the genus Pectinatus

<400> 36  
gtgaagtttt gagtgtgcaa ga 22

<210> 37  
<211> 22  
<212> DNA  
<213> Artifical sequence

<220>  
<223> Description of the artifical sequence: specific  
sequence for the genus Megasphaera

<400> 37  
gaccgaggac ttgacttaag ca 22

<210> 38  
<211> 20  
<212> DNA  
<213> Artifical sequence

|       |  |    |
|-------|--|----|
| <220> |  |    |
| <223> | Description of the artifical sequence: specific sequence for the genus Selenomonas |    |
| <400> | 38   |    |
|       | tccagtgcgc atagctgagt  | 20 |
| <210> | 39   |    |
| <211> | 25   |    |
| <212> | DNA  |    |
| <213> | Artifical sequence   |    |
| <220> |  |    |
| <223> | Description of the artifical sequence: specific sequence for the genus Zymophilus  |    |
| <400> | 39   |    |
|       | agaatatct ggtagtgata gccaa   | 25 |
| <210> | 40   |    |
| <211> | 19   |    |
| <212> | DNA  |    |
| <213> | Artifical sequence   |    |
| <220> |  |    |
| <223> | Description of the artifical sequence: consensus sequence                          |    |
| <400> | 40   |    |
|       | gtcgtgagac agttcggtc   | 19 |
| <210> | 41   |    |
| <211> | 21   |    |
| <212> | DNA  |    |
| <213> | Artifical sequence   |    |
| <220> |  |    |
| <223> | Description of the artifical sequence: consensus sequence                          |    |
| <400> | 41   |    |
|       | cytagtacga gaggaccggr r  | 21 |
| <210> | 42   |    |
| <211> | 21   |    |
| <212> | DNA  |    |
| <213> | Artifical sequence   |    |
| <220> |  |    |
| <223> | Description of the artifical sequence: consensus sequence                          |    |
| <400> | 42   |    |
|       | gctaccctgg ggataacagg c  | 21 |
| <210> | 43   |    |
| <211> | 21   |    |
| <212> | DNA  |    |

<213> Artifical sequence

<220>

<223> Description of the artifical sequence: consensus sequence

<400> 43

atcgacggggg aggtttssca c

21

<210> 44

<211> 20

<212> DNA

<213> Artifical sequence

<220>

<223> Description of the artifical sequence: consensus sequence

<400> 44

cacctcgatg tcggctcrtc

20

<210> 45

<211> 18

<212> DNA

<213> Artifical sequence

<220>

<223> Description of the artifical sequence: consensus sequence

<400> 45

ccaagggttg ggctgttc

18

<210> 46

<211> 19

<212> DNA

<213> Artifical sequence

<220>

<223> Description of the artifical sequence: consensus sequence

<400> 46

aagggccatc rctcaacgg

19

<210> 47

<211> 20

<212> DNA

<213> Artifical sequence

<220>

<223> Description of the artifical sequence: consensus sequence

<400> 47

aagtgtgtaa agcatctaag

20

<210> 48

<211> 23

<212> DNA

<213> Artifical sequence

<220>

<223> Description of the artifical sequence: consensus sequence

<220>

<221> misc\_feature

<222> (9)..(10)

<223> "n" is inosine

<400> 48

tgtgttcgnn atgggaacag gtg

23

<210> 49

<211> 23

<212> DNA

<213> Artifical sequence

<220>

<223> Description of the artifical sequence: consensus sequence

<400> 49

tgtgttcgga atgggaacag gtg

23

<210> 50

<211> 23

<212> DNA

<213> Artifical sequence

<220>

<223> Description of the artifical sequence: consensus sequence

<400> 50

tgtgttcgaa atgggaacag gtg

23

<210> 51

<211> 23

<212> DNA

<213> Artifical sequence

<220>

<223> Description of the artifical sequence: consensus sequence

<400> 51

tgtgttcggt atgggaacag gtg

23

<210> 52

<211> 23

<212> DNA

<213> Artifical sequence

<220>

<223> Description of the artifical sequence: consensus sequence

<400> 52

tgtgttcgat atgggaacag gtg

23

<210> 53  
 <211> 23  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: consensus sequence

<400> 53  
 tgtgttcggc atgggaacag gtg 23

<210> 54  
 <211> 23  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: consensus sequence

<400> 54  
 tgtgttcgac atgggaacag gtg 23

<210> 55  
 <211> 19  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: consensus sequence

<400> 55  
 ggcrrygtcc taytytcsc 19

<210> 56  
 <211> 19  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: consensus sequence

<400> 56  
 ggcagtgtcc tactttccc 19

<210> 57  
 <211> 19  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: consensus sequence

<400> 57  
 ggcagcgtcc tactttcgc 19

```
<210> 58
<211> 19
<212> DNA
<213> Artificial sequence
```

```
<220>
<223> Description of the artifical sequence: consensus sequence
```

```
<400> 58
ggcagtgtcc tacttttcgc
```

```
<210> 59
<211> 19
<212> DNA
<213> Artificial sequence
```

```
<220>
<223> Description of the artifical sequence: consensus sequence
```

```
<400> 59
ggcagcgtcc tactttccc
```

```
<210> 60
<211> 18
<212> DNA
<213> Artificial sequence
```

```
<220>
<223> Description of the artifical sequence: consensus sequence
```

<400> 60  
gyttmrettc yrdgttcg 18

```
<210> 61
<211> 18
<212> DNA
<213> Artifical sequence
```

```
<220>
<223> Description of the artifical sequence: consensus sequence
```

```
<400> 61
gcttaacttc cgtgttcg
```

```
<210> 62
<211> 18
<212> DNA
<213> Artificial sequence
```

<220>  
<223> Description of the artificial sequence: consensus sequence

```
<400> 62
qcttaacttc tatgttcg
```

<210> 63

<211> 18  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: consensus sequence

<400> 63  
 gcttaacttc tgtgttcg 18

<210> 64  
 <211> 18  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: consensus sequence

<400> 64  
 gcttaacttc catgttcg 18

<210> 65  
 <211> 18  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: consensus sequence

<400> 65  
 gcttaacttc cgggttcg 18

<210> 66  
 <211> 18  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: consensus sequence

<400> 66  
 gcttaacttc taggttcg 18

<210> 67  
 <211> 18  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: consensus sequence

<400> 67  
 gcttaacttc tgggttcg 18

<210> 68  
 <211> 18

<212> DNA  
 <213> Artifical sequence

<220>

<223> Description of the artifical sequence: consensus sequence

<400> 68  
 gcttaacttc caggttcg 18

<210> 69  
 <211> 18  
 <212> DNA  
 <213> Artifical sequence

<220>

<223> Description of the artifical sequence: consensus sequence

<400> 69  
 gcttaacttc cgagttcg 18

<210> 70  
 <211> 18  
 <212> DNA  
 <213> Artifical sequence

<220>

<223> Description of the artifical sequence: consensus sequence

<400> 70  
 gcttaacttc taagttcg 18

<210> 71  
 <211> 18  
 <212> DNA  
 <213> Artifical sequence

<220>

<223> Description of the artifical sequence: consensus sequence

<400> 71  
 gcttaacttc tgagttcg 18

<210> 72  
 <211> 18  
 <212> DNA  
 <213> Artifical sequence

<220>

<223> Description of the artifical sequence: consensus sequence

<400> 72  
 gcttaacttc caagttcg 18

<210> 73  
 <211> 25



<212> DNA  
 <213> Artifical sequence  
  
 <220>  
 <223> Description of the artifical sequence: specific  
 sequence for Lactobacillus brevis  
  
 <400> 73  
 tcgagaataa ttgaataata tctag 25  
  
 <210> 74  
 <211> 20  
 <212> DNA  
 <213> Artifical sequence  
  
 <220>  
 <223> Description of the artifical sequence: specific  
 sequence for Lactobacillus brevis  
  
 <400> 74  
 gaggaagaa gttctcttat 20  
  
 <210> 75  
 <211> 23  
 <212> DNA  
 <213> Artifical sequence  
  
 <220>  
 <223> Description of the artifical sequence: specific  
 sequence for Lactobacillus lindneri  
  
 <400> 75  
 aacagagaag atattatcta gtt 23  
  
 <210> 76  
 <211> 42  
 <212> DNA  
 <213> Artifical sequence  
  
 <220>  
 <223> Description of the artifical sequence: specific  
 sequence for Lactobacillus lindneri  
  
 <400> 76  
 ttgagagaac gaagttcgct caggcttatg aaaaataagc at 42  
  
 <210> 77  
 <211> 45  
 <212> DNA  
 <213> Artifical sequence  
  
 <220>  
 <223> Description of the artifical sequence: specific  
 sequence for Lactobacillus casei  
  
 <400> 77  
 ttcgttggcc gggttttggc caatggattc agggttctta tgtgg 45

<210> 78  
 <211> 58  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: specific  
 sequence for Lactobacillus casei

<400> 78  
 gcgtttcgat gaaatacact gggtcccgac aacacaaaaa caacaatgat agccagtt 58

<210> 79  
 <211> 29  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: specific  
 sequence for Lactobacillus casei and Lactobacillus  
 paracasei

<400> 79  
 ttagaaaccg gagcataagc gggcctgag 29

<210> 80  
 <211> 46  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: specific  
 sequence for Lactobacillus paracasei

<400> 80  
 gcgtgatggc cgggcttttg ccattgcggt cagggtcctt atgtgc 46

<210> 81  
 <211> 46  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: specific  
 sequence for Lactobacillus paracasei

<400> 81  
 caagtacggt aagttcaagg cagcaattaa acaatgatag ctagtt 46

<210> 82  
 <211> 44  
 <212> DNA  
 <213> Artifical sequence

<220>

<223> Description of the artifical sequence: specific  
sequence for *Lactobacillus coryniformis*

<400> 82  
aaagaaatga atatccagtt ttgagagcgc aacgttctca gaaa 44

<210> 83  
<211> 48  
<212> DNA  
<213> Artifical sequence

<220>  
<223> Description of the artifical sequence: specific  
sequence for *Lactobacillus curvatus*

<400> 83  
aggtgcaatg ttaggctttt gaaatgaaat attacttatt atgcagtt 48

<210> 84  
<211> 22  
<212> DNA  
<213> Artifical sequence

<220>  
<223> Description of the artifical sequence: specific  
sequence for *Pediococcus damnosus*

<400> 84  
gccgcgtaag tggatcggag aa 22

<210> 85  
<211> 22  
<212> DNA  
<213> Artifical sequence

<220>  
<223> Description of the artifical sequence: specific  
sequence for *Pediococcus inopinatus*

<400> 85  
gccgcggaag tggatcggag aa 22

<210> 86  
<211> 25  
<212> DNA  
<213> Artifical sequence

<220>  
<223> Description of the artifical sequence: sequence for  
the detection of *Pediococcus damnosus*, *Pediococcus*  
*inopinatus* and *Pediococcus parvulus*

<400> 86  
gagagaataa atttctttca cacga 25

<210> 87

<211> 39  
 <212> DNA  
 <213> Artifical sequence

<220>

<223> Description of the artifical sequence: specific  
 sequence for *Pectinatus cerevisiiphilus*

<400> 87  
 aaaatcatcg aaaaaaatgt ttggtctgag attttcttct 39

<210> 88  
 <211> 25  
 <212> DNA  
 <213> Artifical sequence

<220>

<223> Description of the artifical sequence: specific  
 sequence for *Pectinatus cerevisiiphilus*

<400> 88  
 cactctgggtt gaagggcagg gaacg 25

<210> 89  
 <211> 39  
 <212> DNA  
 <213> Artifical sequence

<220>

<223> Description of the artifical sequence: specific  
 sequence for *Pectinatus frisingensis*

<400> 89  
 gatttcatca aaaaagagaa atgtttggtc agagatttt 39

<210> 90  
 <211> 33  
 <212> DNA  
 <213> Artifical sequence

<220>

<223> Description of the artifical sequence: specific  
 sequence for *Pectinatus frisingensis*

<400> 90  
 tatataccgg ctgaggtgct gaggcactga agg 33

<210> 91  
 <211> 36  
 <212> DNA  
 <213> Artifical sequence

<220>

<223> Description of the artifical sequence: specific  
 sequence for *Pectinatus spec. DSM 20764*

<400> 91

aatttcattct ataaatgttt ggtcctgatt tcttct

36

<210> 92  
 <211> 54  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: specific  
 sequence for Pectinatus spec. DSM 20764

<400> 92  
 agattagttc ctggtttact ttatatatga gcactaaggt gcagaaaaga acgt

54

<210> 93  
 <211> 20  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: specific  
 sequence for Pectinatus spec. DSM 20764

<400> 93  
 aggaaacgcg gcgttcgtaa

20

<210> 94  
 <211> 56  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: specific  
 sequence for Selenomonas lacticifex

<400> 94  
 taataatcta gaatgtttcg atacaatttt tcttctgtat agttttgagt ggacat

56

<210> 95  
 <211> 24  
 <212> DNA  
 <213> Artifical sequence

<220>  
 <223> Description of the artifical sequence: specific  
 sequence for Zymophilus raffinivorans

<400> 95  
 gaggcgaaag cggaaggcag cgat

24

<210> 96  
 <211> 24  
 <212> DNA  
 <213> Artifical sequence

<220>

<223> Description of the artifical sequence: specific  
sequence for *Zymophilus paucivorans*

<400> 96  
gaggcgaaag ctaaaggcag cgat

24

<210> 97  
<211> 37  
<212> DNA  
<213> Artifical sequence

<220>  
<223> Description of the artifical sequence: specific  
sequence for *Megasphaera cerevisiae*

<400> 97  
aatcctgaaa cgaattcagt ggtgatggct gcaggga

37

<210> 98  
<211> 20  
<212> DNA  
<213> Artifical sequence

<220>  
<223> Description of the artifical sequence: sequence for  
detection of bacteria of the family Lactobacillaceae  
that are relevant to brewing

<400> 98  
tatggaagta agaccctga

20

<210> 99  
<211> 21  
<212> DNA  
<213> Artifical sequence

<220>  
<223> Description of the artifical sequence: sequence for  
detection of bacteria of the family Lactobacillaceae  
that are relevant to brewing

<400> 99  
agatgatcag gtagataggc t

21

<210> 100  
<211> 21  
<212> DNA  
<213> Artifical sequence

<220>  
<223> Description of the artifical sequence: sequence for  
detection of bacteria of the family Lactobacillaceae  
that are relevant to brewing

<400> 100  
agatgatcag gtcgataggt t

21

<210> 101  
 <211> 21  
 <212> DNA  
 <213> Artifical sequence

<220>

<223> Description of the artifical sequence: sequence for  
 detection of bacteria of the family Lactobacillaceae  
 that are relevant to brewing

<400> 101  
 agatgatcag gtagataggt t

21

<210> 102  
 <211> 25  
 <212> DNA  
 <213> Artifical sequence

<220>

<223> Description of the artifical sequence: sequence for  
 detection of bacteria of the family Lactobacillaceae  
 that are relevant to brewing

<400> 102  
 tactaatcgg tcgaggactt aacca

25

<210> 103  
 <211> 26  
 <212> DNA  
 <213> Artifical sequence

<220>

<223> Description of the artifical sequence: sequence for  
 detection of bacteria of the family Lactobacillaceae  
 that are relevant to brewing

<400> 103  
 atactaatca gtcgaggact taacca

26

<210> 104  
 <211> 32  
 <212> DNA  
 <213> Artifical sequence

<220>

<223> Description of the artifical sequence: specific  
 sequence for the genus Pectinatus

<400> 104  
 gaagcggact ggtactaata agccgagagc tt

32

<210> 105  
 <211> 32  
 <212> DNA  
 <213> Artifical sequence

<220>  
<223> Description of the artifical sequence: specific  
sequence for the genus Selenomonas

<400> 105  
cagcggacca atactaataa atcgagggct ta 32

<210> 106  
<211> 38  
<212> DNA  
<213> Artifical sequence

<220>  
<223> Description of the artifical sequence: specific  
sequence for the genus Zymophilus

<400> 106  
agcggaccga tactaatagg tcgagggctt gacttaaa 38

<210> 107  
<211> 32  
<212> DNA  
<213> Artifical sequence

<220>  
<223> Description of the artifical sequence: specific  
sequence for the genus Megasphaera

<400> 107  
ggagcggacc ggtactaata gaccgaggac tt 32